



STIC Search Report

Biotech-Chem Library

STIC Database Tracking Number: 114839

TO: Jeanine Goldberg
Location: rem/2e74
Art Unit: 1634
Friday, February 20, 2004

Case Serial Number: 10/009897

From: Barb O'Bryen
Location: Biotech-Chem Library
Remsen E01A69
Phone: 571-272-2518 *BOB*

barbara.obryen@uspto.gov

Search Notes

O'Bryen, Barbara

From: Goldberg, Jeanine
Sent: Friday, February 20, 2004 12:17 PM
To: O'Bryen, Barbara
Subject: RE: 10/009897-seq id no: 38

28 Please.
thanks

-----Original Message-----

From: O'Bryen, Barbara
Sent: Friday, February 20, 2004 12:17 PM
To: Goldberg, Jeanine
Subject: RE: 10/009897-seq id no: 38

Jeanine,
In the subject line you have seq 38, but the message has seq 28. Which is correct?
Thanks,
Barb

-----Original Message-----

From: Goldberg, Jeanine
Sent: Friday, February 20, 2004 12:10 PM
To: O'Bryen, Barbara
Subject: 10/009897-seq id no: 38

Can you please tell me if SEQ ID NO: 28 has any hits compared to patent ~~6,495,110~~ searching only against this patent.

THANKS

6492110

Jeanine Enewold Goldberg
1634
571-272-0743
REM 2E74

STIC-ILL

NW

From: Goldberg, Jeanine
Sent: Friday, February 20, 2004 11:46 AM
To: STIC-ILL
Subject: pl ease pull hiv reference

483360

1. Korber et al. 1997 Human Retroviruses and AIDS: a Compilation and analysis of Nucleic Acid and Amino Acid Sequences, Pp. 118-190. Los Alamos, NM Los Alamos National Laboratory..

THANKS

Jeanine Enewold Goldberg
1634
571-272-0743
REM 2E74

506.20

UMM 2/23

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cleic - nucleic search, using sw model

n: February 20, 2004, 14:42:01 ; Search time 4 Seconds
(without alignments)
2.678 Million cell updates/sec

i: ct score: 22

nce: 1 cacaattaaactgtgcattac 22

ng table: IDENTITY NUC

Gapop 10.0 , Gapext 0.5

hed: 112 seqs, 243491 residues

l number of hits satisfying chosen parameters: 224

num DB seq length: 0

num DB seq length: 2000000000

-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

base: 6492110.seq:

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Hit	Score	Query	Length	ID	Description
1	17.4	79.1	2529	1	US-09-184-418C-62
2	17.4	79.1	2601	1	US-09-184-418C-60
3	17.4	79.1	2652	1	US-09-184-418C-61
4	17.4	79.1	8954	1	US-09-184-418C-6
5	15.6	70.9	2574	1	US-09-184-418C-107
6	15.6	70.9	2631	1	US-09-184-418C-104
7	15.6	70.9	2651	1	US-09-184-418C-105
8	15.6	70.9	8959	1	US-09-184-418C-11
9	15.2	69.1	2583	1	US-09-184-418C-44
10	15.2	69.1	2640	1	US-09-184-418C-41
11	15.2	69.1	2658	1	US-09-184-418C-42
12	15.2	69.1	8992	1	US-09-184-418C-4
13	14.2	64.5	2538	1	US-09-184-418C-36
14	14.2	64.5	2538	1	US-09-184-418C-54
15	14.2	64.5	2541	1	US-09-184-418C-18
16	14.2	64.5	2579	1	US-09-184-418C-51
17	14.2	64.5	2606	1	US-09-184-418C-33
18	14.2	64.5	2610	1	US-09-184-418C-15
19	14.2	64.5	2645	1	US-09-184-418C-52
20	14.2	64.5	2657	1	US-09-184-418C-34
21	14.2	64.5	2661	1	US-09-184-418C-16
22	14.2	64.5	8953	1	US-09-184-418C-3
23	14.2	64.5	8966	1	US-09-184-418C-5
24	14.2	64.5	8968	1	US-09-184-418C-1
25	13.6	61.8	2598	1	US-09-184-418C-98
26	13.6	61.8	2655	1	US-09-184-418C-95
27	13.6	61.8	2721	1	US-09-184-418C-96
28	13.6	61.8	9060	1	US-09-184-418C-10
29	13.2	60.0	579	1	US-09-184-418C-75
30	13.2	60.0	579	1	US-09-184-418C-84
31	13.2	60.0	8972	1	US-09-184-418C-9
32	13.2	60.0	9010	1	US-09-184-418C-8
33	12.6	57.3	2544	1	US-09-184-418C-89

ALIGNMENTS

RESULT 1
US-09-184-418C-62/c
; Sequence 62, Application US/09184418C
; Patent No. 6492110
; GENERAL INFORMATION:
; APPLICANT: Hahn, Beatrice
; APPLICANT: Gao, Feng
; APPLICANT: Shaw, George
; TITLE OF INVENTION: CLONES AND SEQUENCES FOR NON-SUBTYPE B ISOLATES OF HUMAN
; TITLE OF INVENTION: IMMUNODEFICIENCY VIRUS TYPE 1
; FILE REFERENCE: D6287
; CURRENT APPLICATION NUMBER: US/09/184,418C
; CURRENT FILING DATE: 1999-11-02
; NUMBER OF SEQ ID NOS: 112
; SEQ ID NO 62
; LENGTH: 2529
; TYPE: DNA
; ORGANISM: Human immunodeficiency virus type 1
; FEATURE:
; OTHER INFORMATION: isolate=93BR029; gene=env
US-09-184-418C-62

Query Match 79.1%; Score 17.4; DB 1; Length 2529;
Best Local Similarity 94.7%; Pred.No.0.33;
Matches 18; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 2 ACAATTAAACTGTGCATT 20

Db 1134 ACAATTAAACTGTGCATT 1116

RESULT 2

US-09-184-418C-60/c
; Sequence 60, Application US/09184418C
; Patent No. 6492110
; GENERAL INFORMATION:
; APPLICANT: Hahn, Beatrice
; APPLICANT: Gao, Feng
; APPLICANT: Shaw, George
; TITLE OF INVENTION: CLONES AND SEQUENCES FOR NON-SUBTYPE B ISOLATES OF HUMAN
; TITLE OF INVENTION: IMMUNODEFICIENCY VIRUS TYPE 1
; FILE REFERENCE: D6287
; CURRENT APPLICATION NUMBER: US/09/184,418C
; CURRENT FILING DATE: 1999-11-02
; NUMBER OF SEQ ID NOS: 112
; SEQ ID NO 60
; LENGTH: 2601
; TYPE: DNA
; ORGANISM: Human immunodeficiency virus type 1
; FEATURE:
; OTHER INFORMATION: isolate=93BR029; gene=tat
US-09-184-418C-60

Query Match 79.1%; Score 17.4; DB 1; Length 2601;
Best Local Similarity 94.7%; Pred.No.0.33;
Matches 18; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

```

: APPLICANT: Gao, Feng
: APPLICANT: Shaw, George
: TITLE OF INVENTION: CLONES AND SEQUENCES FOR NON-SUBTYPE B ISOLATES OF HUMAN
: TITLE OF INVENTION: IMMUNODEFICIENCY VIRUS TYPE 1
: FILE REFERENCE: D6287
: CURRENT APPLICATION NUMBER: US/09/184,418C
: CURRENT FILING DATE: 1999-11-02
:

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SULT 5
-09-164-418C-107/c

NUMBER OF SEQ ID NOS: 112

EQ ID NO 105

LENGTH: 2651

TYPE: DNA

ORGANISM: Human immunodeficiency virus type 1

FEATURE:

OTHER INFORMATION: isolate=94IN476.104; gene=rev

09-184-418C-105

Query Match 70.9%; Score 15.6; DB 1; Length 2651;

est Local Similarity 81.8%; Pred. No. 1.6;

atches 18; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

1 CACAATTAAACTGTGCTTAC 22

|||||

1387 CACAATTAAAGCTATGTGTTAC 1366

ULT 8

09-184-418C-11/c

sequence 11, Application US/09184418C

atent No. 6492110

GENERAL INFORMATION:

APPLICANT: Hahn, Beatrice

APPLICANT: Gao, Feng

APPLICANT: Shaw, George

TITLE OF INVENTION: CLONES AND SEQUENCES FOR NON-SUBTYPE B ISOLATES OF HUMAN

FILE REFERENCE: D6287

CURRENT APPLICATION NUMBER: US/09/184,418C

CURRENT FILING DATE: 1999-11-02

NUMBER OF SEQ ID NOS: 112

EQ ID NO 11

LENGTH: 8959

TYPE: DNA

ORGANISM: Human immunodeficiency virus type 1

FEATURE:

OTHER INFORMATION: isolate=94IN476.104; 138.1613:"gag";

OTHER INFORMATION: 1418.4428:"pol"; 4361.4939:"vif"; 4879.5169:"vpr";

OTHER INFORMATION: 5150.7782:"tat"; 5289.7939:"rev"; 5378.5638:"vpu";

OTHER INFORMATION: 5556.8129:"env"; 8131.8754:"nef

09-184-418C-11

Query Match

est Local Similarity 70.9%; Score 15.6; DB 1; Length 8959;

atches 18; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

1 CACAATTAAACTGTGCTTAC 22

|||||

6675 CACAATTAAAGCTATGTGTTAC 6654

ULT 9

09-184-418C-44/c

sequence 44, Application US/09184418C

atent No. 6492110

GENERAL INFORMATION:

APPLICANT: Hahn, Beatrice

APPLICANT: Gao, Feng

APPLICANT: Shaw, George

TITLE OF INVENTION: CLONES AND SEQUENCES FOR NON-SUBTYPE B ISOLATES OF HUMAN

FILE REFERENCE: D6287

CURRENT APPLICATION NUMBER: US/09/184,418C

CURRENT FILING DATE: 1999-11-02

NUMBER OF SEQ ID NOS: 112

EQ ID NO 44

LENGTH: 2583

TYPE: DNA

ORGANISM: Human immunodeficiency virus type 1

FEATURE:

OTHER INFORMATION: isolate=92RW009; gene=env

09-184-418C-44

Query Match 69.1%; Score 15.2; DB 1; Length 2583;

Best Local Similarity 85.0%; Pred. No. 2.3;

Matches 17; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 CACAATTAAACTGTGCTTAC 20

|||||

DB 1135 CACAATTAAACTATGTGTT 1116

RESULT 10

US-09-184-418C-41/c

; Sequence 41, Application US/09184418C

; Patent No. 6492110

; GENERAL INFORMATION:

; APPLICANT: Hahn, Beatrice

; APPLICANT: Gao, Feng

; APPLICANT: Shaw, George

; TITLE OF INVENTION: CLONES AND SEQUENCES FOR NON-SUBTYPE B ISOLATES OF HUMAN

; TITLE OF INVENTION: IMMUNODEFICIENCY VIRUS TYPE 1

; FILE REFERENCE: D6287

; CURRENT APPLICATION NUMBER: US/09/184,418C

; CURRENT FILING DATE: 1999-11-02

; NUMBER OF SEQ ID NOS: 112

; SEQ ID NO 41

; LENGTH: 2640

; TYPE: DNA

; ORGANISM: Human immunodeficiency virus type 1

; FEATURE:

; OTHER INFORMATION: isolate=92RW009; gene=tat

US-09-184-418C-41

Query Match

Best Local Similarity 69.1%; Score 15.2; DB 1; Length 2640;

Matches 17; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 CACAATTAAACTGTGCTTAC 20

|||||

DB 1539 CACAATTAAACTATGTGTT 1520

RESULT 11

US-09-184-418C-42/c

; Sequence 42, Application US/09184418C

; Patent No. 6492110

; GENERAL INFORMATION:

; APPLICANT: Hahn, Beatrice

; APPLICANT: Gao, Feng

; APPLICANT: Shaw, George

; TITLE OF INVENTION: CLONES AND SEQUENCES FOR NON-SUBTYPE B ISOLATES OF HUMAN

; TITLE OF INVENTION: IMMUNODEFICIENCY VIRUS TYPE 1

; FILE REFERENCE: D6287

; CURRENT APPLICATION NUMBER: US/09/184,418C

; CURRENT FILING DATE: 1999-11-02

; NUMBER OF SEQ ID NOS: 112

; SEQ ID NO 42

; LENGTH: 2658

; TYPE: DNA

; ORGANISM: Human immunodeficiency virus type 1

; FEATURE:

; OTHER INFORMATION: isolate=92RW009; gene=rev

US-09-184-418C-42

Query Match

Best Local Similarity 69.1%; Score 15.2; DB 1; Length 2658;

Matches 17; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 CACAATTAAACTGTGCTTAC 20

|||||

DB 1400 CACAATTAAACTATGTGTT 1381

RESULT 12

```
3-09-184-418C-4/c
Sequence 4, Application US/09184418C
Patent No. 6492110
GENERAL INFORMATION:
APPLICANT: Hahn, Beatrice
APPLICANT: Gao, Feng
APPLICANT: Shaw, George
TITLE OF INVENTION: CLONES AND SEQUENCES FOR NON-SUBTYPE B ISOLATES OF HUMAN
TITLE OF INVENTION: IMMUNODEFICIENCY VIRUS TYPE 1
FILE REFERENCE: D6287
CURRENT APPLICATION NUMBER: US/09/184.418C
CURRENT FILING DATE: 1999-11-02
NUMBER OF SEQ ID NOS: 112
SEQ ID NO 4
LENGTH: 8992
TYPE: DNA
ORGANISM: Human immunodeficiency virus type 1
FEATURE:
OTHER INFORMATION: isolate=92RM009; 139.1624:gag; 1690.4428:pol (N-terminus uncertain)
OTHER INFORMATION: 4373.4951:vif; 4891.5181:vpr; 5162.7801:tat; 5301.7958:rev;
OTHER INFORMATION: 5403.5648:vpu; 5566.8148:env; 8150.8773:nef
3-09-184-418C-4
Query Match 69.1%; Score 15.2; DB 1; Length 8992;
Best Local Similarity 85.0%; Pred. No. 2.9;
Matches 17; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

1 CACAATTAATAACTGTGCATT 20
|||||
6700 CACAATTAATAACTGTGTT 6681

RESULT 13
3-09-184-418C-36/c
Sequence 36, Application US/09184418C
Patent No. 6492110
GENERAL INFORMATION:
APPLICANT: Hahn, Beatrice
APPLICANT: Gao, Feng
APPLICANT: Shaw, George
TITLE OF INVENTION: CLONES AND SEQUENCES FOR NON-SUBTYPE B ISOLATES OF HUMAN
TITLE OF INVENTION: IMMUNODEFICIENCY VIRUS TYPE 1
FILE REFERENCE: D6287
CURRENT APPLICATION NUMBER: US/09/184.418C
CURRENT FILING DATE: 1999-11-02
NUMBER OF SEQ ID NOS: 112
SEQ ID NO 36
LENGTH: 2538
TYPE: DNA
ORGANISM: Human immunodeficiency virus type 1
FEATURE:
OTHER INFORMATION: isolate=90CR056; gene=env
3-09-184-418C-36
Query Match 64.5%; Score 14.2; DB 1; Length 2538;
Best Local Similarity 84.2%; Pred. No. 5.4;
Matches 16; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

2 ACAATTAATAACTGTGCATT 20
|||||
1107 ACAATTAATAACTGTGTT 1089

RESULT 14
3-09-184-418C-54/c
Sequence 54, Application US/09184418C
Patent No. 6492110
GENERAL INFORMATION:
APPLICANT: Hahn, Beatrice
APPLICANT: Gao, Feng
APPLICANT: Shaw, George
TITLE OF INVENTION: CLONES AND SEQUENCES FOR NON-SUBTYPE B ISOLATES OF HUMAN
TITLE OF INVENTION: IMMUNODEFICIENCY VIRUS TYPE 1
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FILE REFERENCE: D6287
CURRENT APPLICATION NUMBER: US/09/184.418C
CURRENT FILING DATE: 1999-11-02
NUMBER OF SEQ ID NOS: 112
SEQ ID NO 54
LENGTH: 2538
TYPE: DNA
ORGANISM: Human immunodeficiency virus type 1
FEATURE:
OTHER INFORMATION: isolate=92NG003; gene=env
US-09-184-418C-54
Query Match 64.5%; Score 14.2; DB 1; Length 2538;
Best Local Similarity 84.2%; Pred. No. 5.4;
Matches 16; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

2 ACAATTAATAACTGTGCATT 20
|||||
1119 ACAATTAATAACTGTGTT 1101

RESULT 15
US-09-184-418C-18/c
Sequence 18, Application US/09184418C
Patent No. 6492110
GENERAL INFORMATION:
APPLICANT: Hahn, Beatrice
APPLICANT: Gao, Feng
APPLICANT: Shaw, George
TITLE OF INVENTION: CLONES AND SEQUENCES FOR NON-SUBTYPE B ISOLATES OF HUMAN
TITLE OF INVENTION: IMMUNODEFICIENCY VIRUS TYPE 1
FILE REFERENCE: D6287
CURRENT APPLICATION NUMBER: US/09/184.418C
CURRENT FILING DATE: 1999-11-02
NUMBER OF SEQ ID NOS: 112
SEQ ID NO 18
LENGTH: 2541
TYPE: DNA
ORGANISM: Human immunodeficiency virus type 1
FEATURE:
OTHER INFORMATION: isolate=93BR020; gene=env
US-09-184-418C-18
Query Match 64.5%; Score 14.2; DB 1; Length 2541;
Best Local Similarity 84.2%; Pred. No. 5.4;
Matches 16; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

2 ACAATTAATAACTGTGCATT 20
|||||
1134 ACAATTAATAACTGTGCTT 1116

Search completed: February 20, 2004, 14:42:06
Job time : 5 secs
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